Figure 1

GTCGTTAAMGGGACCTGGGAAGGAGCATAGGACAAGGCCAAGGCGGGATAAGGAGGGGGCACCACACACGTTAAGGCACGAGGGAAA

Xagext.4

Are CTC CTT TGG TGC CCA CCT CAG TGC GCA TGT TCA CTG GGC GTC TTC CCA 147/17. Met leu l'eu trp cys pro pro gln cys ala cys ser leu gly val phe pro CCTCACTGCGC ATG CTC

207/37 GCC CCT TCG CCA GTG TGG GGA ACG CGG AGC TGT GAG CCG GCG ACT CGG GTC CCT ala pro ser pro val trp gly thr arg arg ser cys glu pro ala thr arg val pro rc ser

GAG GTC TGG ATT CTT TCT CCG CTA CTG AGA CAC GGC GCA CAC ACA CAA ACA CAG AAC CAC 267/57 glu val trp ile leu ser pro leu leu arg his gly gly his thr gln thr gln asn his Xagext.3

Xa-2

ACA GCC AGT CCC AGG AGC CCA GTA ATG GAG AGC CCC AAA AAG AAG AAC CAG CAG CTG AAA 327/77 thr ala ser pro arg ser pro val Mat glu ser pro lys lys asn gln gln leu lys

GTC GGG ATC CTA CAC CTG GGC AGC AGA CAG AAG AAG ATC AGG ATA CAG CTG AGA TCC CAG 387/97 val gly ile leu his leu gly ser arg gln lys lys ile arg ile gln leu arg ser gln

TGC GCG ACA TGG AAG GTG ATC TGC AAG AGC TGC ATC AGT CAA ACA CCG GGG ATA AAT CTG 447/117 cys ala thr trp lys val ile cys lys ser cys ile ser gln thr pro gly ile asn leu

GAT TTG GGT TCC GGC GTC AAG GTG AAG ATA ATA CCT AAA GAG GAA CAC TGT AAA ATG CCA 507/137 asp leu gly ser gly val lys val lys lie lie pro lys glu glu his cys lys met pro

gaa gca d<mark>o</mark>t gaa gag caa cca caa git taa atgaagacaagctgaaacaaggcaagctggtttaatattagatat glu ala gly glu glu gln pro gln val och

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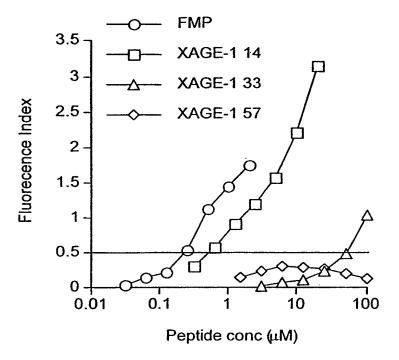


Figure 2

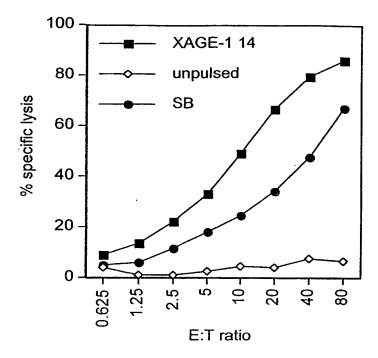


Figure 3